SEQUENCE LISTING



RECEIVED

JUL 2 7 2001

TECH CENTER 1600/2900

<110> DUNICAN, L.K.
 MCCORMACK, ASHLING
 STAPELTON, CLIONA
 BURKE, KEVIN
 MOCKEL, BETTINA

<120> NEW NUCLEOTIDE SEQUENCES WHICH CODE FOR THE TAL GENE

<130> MAS/21123/258100

<140> 09/531,266

<141> 2000-03-20

<150> 60/142,915

<151> 1999-07-09

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<170> PatentIn Ver. 2.1

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				tcc Ser			His										3085
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			cag Gln													3469
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taaa	agaaa	agg a	atcgi	gaca	ac ta	accat	tcgtg	g ago	cacaa	aaca	cgad	cccc	etc o	cagct	ggaca	3690
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Ala Ala Met Ser Lys Gly Asp Ser Tyr Asp Ala Gln Ile Ala Glu 50 55 60

Leu Lys Ala Ala Gly Ala Ser Val Asp Gln Ala Val Tyr Ala Met Ser 65 70 75 80

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Asp Asp Leu Ser Arg Glu Arg Ile Thr Ser Gly Asn Leu Ser Gln Val
att gag gaa aag tot gta gto ggt gto acc acc ca got att tto
Ile Glu Glu Lys Ser Val Val Gly Val Thr Thr Asn Pro Ala Ile Phe
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Ala Ala Met Ser Lys Gly Asp Ser Tyr Asp Ala Gln Ile Ala Glu
ctc aag gcc gct ggc gca tct gtt gac cag gct gtt tac gcc atg agc
Leu Lys Ala Ala Gly Ala Ser Val Asp Gln Ala Val Tyr Ala Met Ser
ate gae gat ege aat get tgt gat etg tte ace gge ate tte gag
                                                                   288
Ile Asp Asp Val Arg Asn Ala Cys Asp Leu Phe Thr Gly Ile Phe Glu
tee tee aac gge tac gac gge ege gtg tee ate gag gtt gac eea egt
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Ser Ser Asn Gly Tyr Asp Gly Arg Val Ser Ile Glu Val Asp Pro Arg
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Ile Ser Ala Asp Arg Asp Ala Thr Leu Ala Gln Ala Lys Glu Leu Trp
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gca aag gtt gat cgt cca aac gtc atg atc aag atc cct gca acc cca
                                                                   432
Ala Lys Val Asp Arg Pro Asn Val Met Ile Lys Ile Pro Ala Thr Pro
                        135
ggt tet ttg cca gca atc acc gac get ttg get gag ggc atc agc gtt
                                                                   480
Gly Ser Leu Pro Ala Ile Thr Asp Ala Leu Ala Glu Gly Ile Ser Val
                    150
                                        155
aac gtc acc ttg atc ttc tcc gtt gct cgc tac cgc gag gtc atc gct
                                                                   528
Asn Val Thr Leu Ile Phe Ser Val Ala Arg Tyr Arg Glu Val Ile Ala
geg tte ate gag gge ate aag eag get get gea aac gge eac gae gte
Ala Phe Ile Glu Gly Ile Lys Gln Ala Ala Ala Asn Gly His Asp Val
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gag Glu	atc Ile 210	gac Asp	aag Lys	cgc Arg	ctc Leu	gag Glu 215	gca Ala	atc Ile	gga Gly	tcc Ser	gat Asp 220	gag Glu	gct Ala	ttg Leu	gct Ala	672
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<212> PRT

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Ile Glu Glu Lys Ser Val Val Gly Val Thr Thr Asn Pro Ala Ile Phe Ala Ala Ala Met Ser Lys Gly Asp Ser Tyr Asp Ala Gln Ile Ala Glu Leu Lys Ala Ala Gly Ala Ser Val Asp Gln Ala Val Tyr Ala Met Ser Ile Asp Asp Val Arg Asn Ala Cys Asp Leu Phe Thr Gly Ile Phe Glu Ser Ser Asn Gly Tyr Asp Gly Arg Val Ser Ile Glu Val Asp Pro Arg 105 Ile Ser Ala Asp Arg Asp Ala Thr Leu Ala Gln Ala Lys Glu Leu Trp 120 Ala Lys Val Asp Arg Pro Asn Val Met Ile Lys Ile Pro Ala Thr Pro 135 Gly Ser Leu Pro Ala Ile Thr Asp Ala Leu Ala Glu Gly Ile Ser Val 150 155 Asn Val Thr Leu Ile Phe Ser Val Ala Arg Tyr Arg Glu Val Ile Ala 170 Ala Phe Ile Glu Gly Ile Lys Gln Ala Ala Ala Asn Gly His Asp Val 180 Ser Lys Ile His Ser Val Ala Ser Phe Phe Val Ser Arg Val Asp Val 200 Glu Ile Asp Lys Arg Leu Glu Ala Ile Gly Ser Asp Glu Ala Leu Ala 210 Leu Arg Gly Lys Ala Gly Val Ala Asn Ala Gln Arg Ala Tyr Ala Val 230 235 Tyr Lys Glu Leu Phe Asp Ala Ala Glu Leu Pro Glu Gly Ala Asn Thr 245

245 250 255

Gln Arg Pro Leu Trp Ala Ser Thr Gly Val Lys Asn Pro Ala Tyr Ala 260 265 270

Ala Thr Leu Tyr Val Ser Glu Leu Ala Gly Pro Asn Thr Val Asn Thr 275 280 285

Met Pro Glu Gly Thr Ile Asp Ala Val Leu Glu Gln Gly Asn Leu His 290 295 300

Gly Asp Thr Leu Ser Asn Ser Ala Ala Glu Ala Asp Ala Val Phe Ser 305 310 315 320

Gln Leu Glu Ala Leu Gly Val Asp Leu Ala Asp Val Phe Gln Val Leu 325 330 335

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